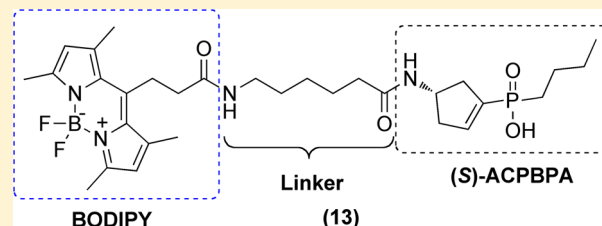


Design, Synthesis, and Pharmacological Evaluation of Fluorescent and Biotinylated Antagonists of ρ_1 GABA_C ReceptorsNavnath Gavande,[†] Hye-Lim Kim,[†] Munikumar R. Doddareddy,[†] Graham A. R. Johnston,[‡] Mary Chebib,[†] and Jane R. Hanrahan^{*†}[†]Faculty of Pharmacy, The University of Sydney, Sydney, NSW 2006, Australia[‡]Adrien Albert Laboratory, Department of Pharmacology, The University of Sydney, Sydney, NSW 2006, Australia

Supporting Information

ABSTRACT: The ρ_1 GABA_C receptor is a ligand-gated chloride ion channel that shows promise as a therapeutic target for myopia, sleep disorders, memory and learning facilitation, and anxiety-related disorders. As such, there is a need for molecular probes to understand the role GABA_C receptors play in physiological and pathological processes. To date, no labeled (either radioactive or fluorescent) GABA_C selective ligand has been developed that can act as a marker for GABA_C receptor visualization and localization studies. Herein, we report a series of fluorescent ligands containing different-sized linkers and fluorophores based around (S)-4-ACPBPA [(4-aminocyclopenten-1-yl)-butylphosphinic acid], a selective GABA_C antagonist. One of these conjugates, (S)-4-ACPBPA-CS-BODIPY (13), displayed moderate potency (IC₅₀ = 58.61 μ M) and selectivity (>100 times) for ρ_1 over $\alpha_1\beta_2\gamma_{2L}$ GABA_A receptors. These conjugates are novel lead agents for the development of more potent and selective fluorescent probes for studying the localization and function of GABA_C receptors in living cells.

KEYWORDS: Human ρ_1 GABA_C receptors, CNS-related disorders, GABA_C antagonists, fluorescent and biotinylated probes, homology modeling and docking



γ -Aminobutyric acid **1** [GABA (Figure 1)] is the most abundant inhibitory neurotransmitter in the mammalian central nervous system (CNS) and is essential for the overall balance between neuronal excitation and inhibition.^{1,2} GABA released from GABAergic axon terminals influences neurons via GABA_A, GABA_B, and GABA_C (GABA ρ) receptors. These receptors are grouped on the basis of their subunit composition, gating properties, and pharmacological profiles. GABA_A and GABA_C receptors are ligand-gated chloride ion channels that mediate fast synaptic inhibition when activated by GABA,¹ whereas GABA_B receptors are G-protein-coupled receptors that mediate slow, longer-lasting synaptic inhibition by increasing potassium and decreasing calcium conductances when activated by GABA.³

Ionotropic GABA_A receptors are transmembrane protein complexes composed of five heteropentameric subunits. To date, 16 human GABA_A receptor subunits have been identified and classified as α (α_1 – α_6), β (β_1 – β_3), γ (γ_1 – γ_3), δ , ϵ , π , and θ .⁴ In contrast, ionotropic GABA_C receptors have pharmacology, physiology, and subunit compositions distinct from those of GABA_A.^{5,6} In mammals, GABA_C receptors are composed of ρ subunits (ρ_1 – ρ_3), forming homopentameric assemblies of five ρ subunits (ρ_1 , ρ_2 , or ρ_3 subunits) or pseudoheteropentameric complexes comprising different ρ subunits (a combination of ρ_1 and ρ_2 or ρ_2 and ρ_3 subunits).^{5,7,8} GABA_C receptors are insensitive to bicuculline and (–)-baclofen, selectively activated by (+)-CAMP **3** [(+)-*cis*-2-(aminomethyl)-

cyclopropanecarboxylic acid (Figure 1)] and 5-Me-I-4AA **4** (5-methyl-1*H*-imidazole-4-acetic acid), and inhibited by TPMPA **5** (1,2,5,6-tetrahydropyridine-4-yl-methyl-phosphinic acid), (S)-4-ACPBPA **6** [(S)-4-aminocyclopenten-1-yl]-butylphosphinic acid⁹ and 3-GOHP **8** [3-(guanido)-1-oxo-1-hydroxy-phospholane].¹⁰ This clearly indicates that the GABA binding sites of GABA_C and GABA_A receptors are not identical.

GABA_A receptors are widely distributed in the CNS,¹ whereas GABA_C receptors are mainly expressed in the superior colliculus,¹¹ cerebellum,¹² lateral amygdala,¹³ hippocampus (strong ρ_2 subunit expression),¹⁴ and, most prominently, the retina (strong ρ_1 subunit expression).⁷ Various studies show that GABA_C receptors are potential therapeutic targets for myopia, sleep disorders, managing memory-related disorders, and peripheral antinociception.^{5,13} However, understanding the role of GABA_C receptors and information about the processes triggered by ligand–receptor interactions in living cells are still limited because of the lack of pharmacological tools.

Fluorescent ligands have proven to be useful tools offering a wealth of information about the mapping or identification of ligand binding sites,^{15,16} the mechanism of ligand binding,¹⁷ the physical nature of the binding pocket, the movement and

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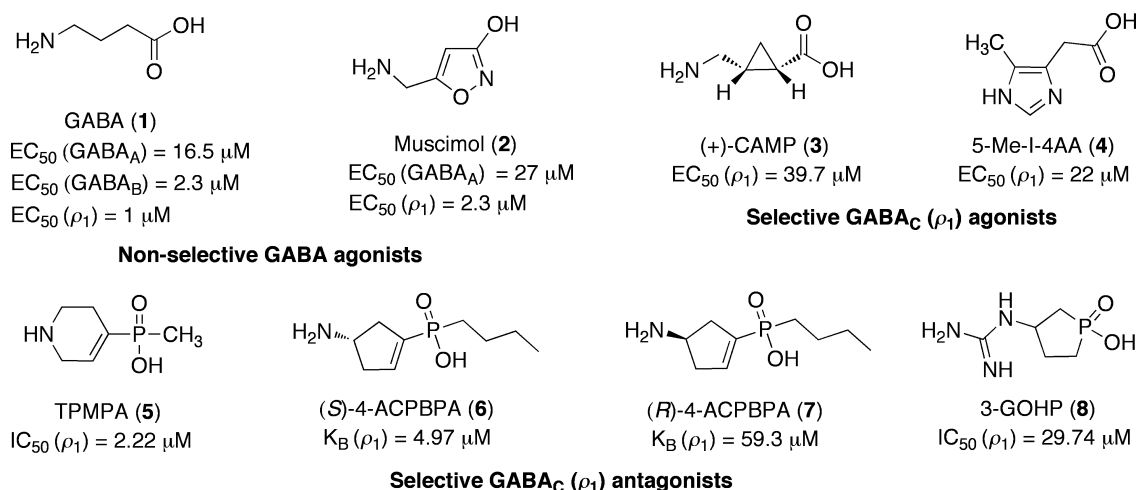
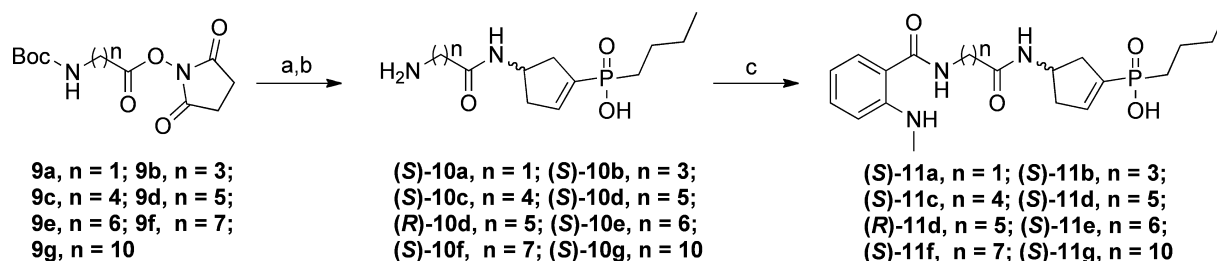


Figure 1. Structures of GABA (1), the GABA_A agonist and GABA_C partial agonist muscimol (2), the GABA_C selective agonists (+)-CAMP (3) and 5-Me-I-4AA (4), and GABA_C selective antagonists TPMPA (5), (S)-4-ACPBPA (6), (R)-4-ACPBPA (7), and 3-GOHP (8).

Scheme 1. Synthesis of Fluorescent ρ_1 GABA_C Antagonists (S)-11a–g and (R)-11d^a



^aReagents and conditions: (a) (S)-4-ACPBPA 6 or (R)-4-ACPBPA 7 [for (R)-10d], NaHCO₃, water/DME/THF (1:1:0.3), room temperature for 16 h, 81–89%; (b) TFA/DCM (1:1), room temperature for 1–2 h, 92–98%; (c) NMA *N*-succinimidyl ester, NaHCO₃, water/DME/THF (1:1:0.3), room temperature for 16 h, 75–84%.

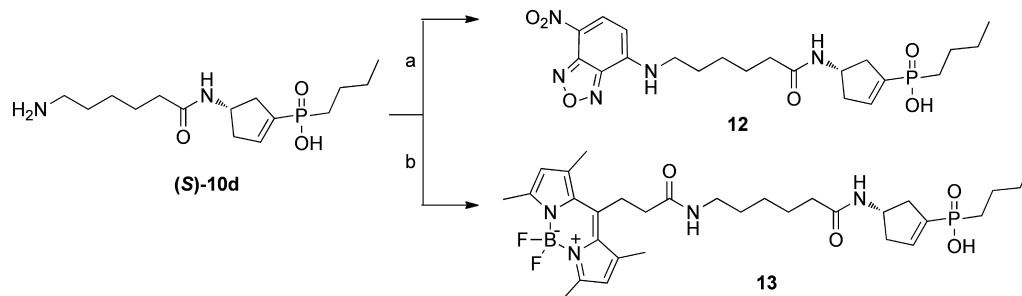
internalization of receptors in living cells,¹⁸ and the localization as well as visualization of the labeled receptors.¹⁹ Biotinylated probes are useful for the isolation and purification of target proteins of the receptor fragment for crystallization and X-ray studies of the particular receptor-binding site,²⁰ the localization and visualization of the receptor by using antibodies,²¹ and the study of ligand–protein interactions in living cells.²²

In addition, fluorescent and biotinylated probes represent a faster, safer, and less expensive alternative to radioligands in receptor studies, circumventing several drawbacks associated with radioligand studies such as health and safety concerns and the need for a large number of cells with strong receptor expression.²³ Thus, the identification of fluorescent or biotinylated ligands for GABA_C receptors is of great interest. The development of these tagged ligands usually starts with the selection of a potent and selective pharmacophore, which is then conjugated with the desired tag through an appropriate linker.

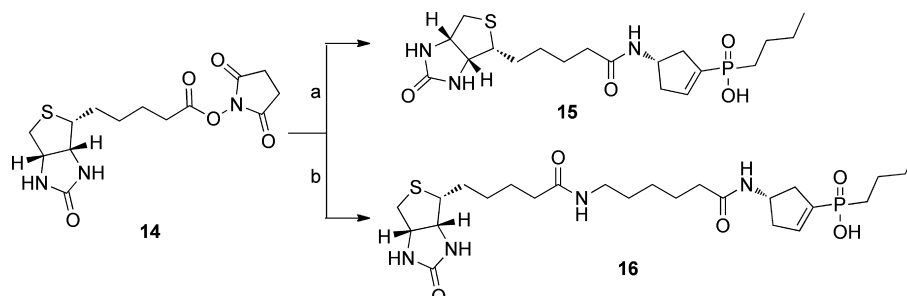
To date, only a few fluorescent and biotinylated GABA_B receptor antagonist ligands have been reported,²⁴ and *N*-acylation of the amino moiety of muscimol (2), a widely used GABA_A agonist that it is also a potent partial agonist at human ρ_1 GABA_C receptors with fluorophores, has been shown to result in ligands that bind to the GABA_A receptors.²⁵ Vu et al. presented evidence that muscimol linked through a 6-aminohexanoyl chain to biotin (muscimol-biotin; EC_{50} = 20 μ M) and BODIPY (muscimol-BODIPY) retained substantial agonist activity compared to muscimol (2; EC_{50} = 2 μ M) at

GABA_C receptors,²⁶ indicating a steric tolerance in the region around the ligand's amine binding site of the GABA_C receptor. To the best of our knowledge, no selective fluorescent ligand for GABA_C receptors has been described in the literature. Therefore, we envisioned that incorporating fluorophores or the versatile biotin moiety into selective GABA_C ligands could lead to selective probes. Herein, we report the design, synthesis, and pharmacological evaluation of the selective fluorescent and biotinylated probes for ρ_1 GABA_C receptors.

We selected two of our previously reported selective and potent GABA_C antagonists, (S)-4-ACPBPA 6 and (R)-4-ACPBPA 7 (Figure 1), for the design of the fluorescent GABA_C antagonists.⁹ Both were synthesized according to our previously published protocol.⁹ The length of the spacer chain is known to significantly impact the affinity of ligand–fluorophore conjugates for the receptors; therefore, we chose a linker size that varied from 0 to 10 carbons to introduce molecular flexibility into the conjugated fluorescent ligand. We incorporated fluorescent groups with relatively small to large molecular volumes to study the effect of perturbation on receptor affinity. We used the fluorophores NMA (*N*-methylanthranilic acid), BODIPY (4,4-difluoro-4-bora-3a,4a-diaza-*s*-indacene), and NBD (7-nitrobenz-2-oxa-1,3-diazol-4-yl) chloride, which are known to have different excitation wavelengths and distinctive properties, to diversify fluorescence experiments. BODIPY has been extensively utilized for the development of fluorescent probes because of its distinctive and useful features, including hydrophobicity, photochemical

Scheme 2. Synthesis of Fluorescent ρ_1 GABA_C Antagonists 12 and 13^a

^aReagents and conditions: (a) 7-nitrobenz-2-oxa-1,3-diazol-4-yl chloride (NBD-Cl), K₂CO₃, DMF, room temperature for 24 h, 76%; (b) BODIPY N-succinimidyl ester, DIPEA, DMF, room temperature for 24 h, 72%.

Scheme 3. Synthesis of Biotinylated ρ_1 GABA_C Antagonists 15 and 16^a

^aReagents and conditions: (a) (S)-4-ACBPBA 6, DIPEA, DMF, room temperature for 24 h, 75%; (b) (S)-10d, DIPEA, DMF, room temperature for 24 h, 81%.

Table 1. Pharmacological Evaluation of Fluorescent and Biotinylated Probes^a

compound	spacer	human ρ_1 GABA _C receptor % inhibition and IC ₅₀ μ M (95% confidence interval)		human $\alpha_1\beta_2\gamma_{2L}$ GABA _A receptor % inhibition at 300 μ M
		300 μ M	600 μ M ^d	
(S)-4-ACBPBA (6)		97 \pm 3% ^b IC ₅₀ = 9.76, K _B = 4.97 μ M ^c		inactive ^g
(R)-4-ACBPBA (7)		92 \pm 7% ^b K _B = 59.3 μ M ^c		inactive ^g
(S)-4-ACBPBA-C1-NMA [(S)-11a]	1		8 \pm 2%	
(S)-4-ACBPBA-C3-NMA [(S)-11b]	3		23 \pm 4%	
(S)-4-ACBPBA-C4-NMA [(S)-11c]	4		31 \pm 2%	
(S)-4-ACBPBA-C5-NMA [(S)-11d]	5		62 \pm 7%	inactive ^g
(R)-4-ACBPBA-C5-NMA [(R)-11d]	5		56 \pm 9%	inactive ^g
(S)-4-ACBPBA-C6-NMA [(S)-11e]	6		55 \pm 3%	
(S)-4-ACBPBA-C7-NMA [(S)-11f]	7		24 \pm 8%	
(S)-4-ACBPBA-C10-NMA [(S)-11g]	10		8 \pm 3%	
(S)-4-ACBPBA-C5-NBD (12)	5	IC ₅₀ = 281.42 (274.96–332.11) ^e IC ₅₀ = 193.27 (181.67–224.81) ^f		inactive ^g
(S)-4-ACBPBA-C5-BODIPY (13)	5	IC ₅₀ = 103.14 (95.36–114.87) ^e IC ₅₀ = 58.61 (45.37–71.63) ^f		inactive ^g
(S)-4-ACBPBA-C0-biotin (15)	0	inactive ^b		
(S)-4-ACBPBA-C5-biotin (16)	5	IC ₅₀ = 147.92 (139.45–174.13) ^e IC ₅₀ = 76.54 (68.33–102.96) ^f		9 \pm 7% ^g

^aDetermined electrophysiologically in *Xenopus laevis* oocytes expressing the human ρ_1 GABA_C receptor or human $\alpha_1\beta_2\gamma_{2L}$ GABA_A receptors as previously described.^{9,10} ^bActivity or percent inhibition by 300 μ M compound of the current produced by a submaximal concentration of GABA (1 μ M, EC₅₀). ^cData from ref 9. ^dPercent inhibition by 600 μ M compound of the current produced by a submaximal concentration of GABA (1 μ M, EC₅₀), with a 5 min preincubation. ^eIC₅₀ values for compounds without preincubation. ^fIC₅₀ values for compounds with a 5 min preincubation. ^gActivity or percent inhibition by 300 μ M compound of the current produced by a submaximal concentration of GABA (30 μ M, EC₅₀) without and with a 5 min preincubation. All data are means \pm the standard error of the mean (n = 3 oocytes).

stability, insensitivity to changes in experimental conditions, efficient uptake into cell membranes, and high extinction

coefficient and fluorescence quantum yield.²⁷ Initially, we synthesized fluorescent probes using the potent and selective

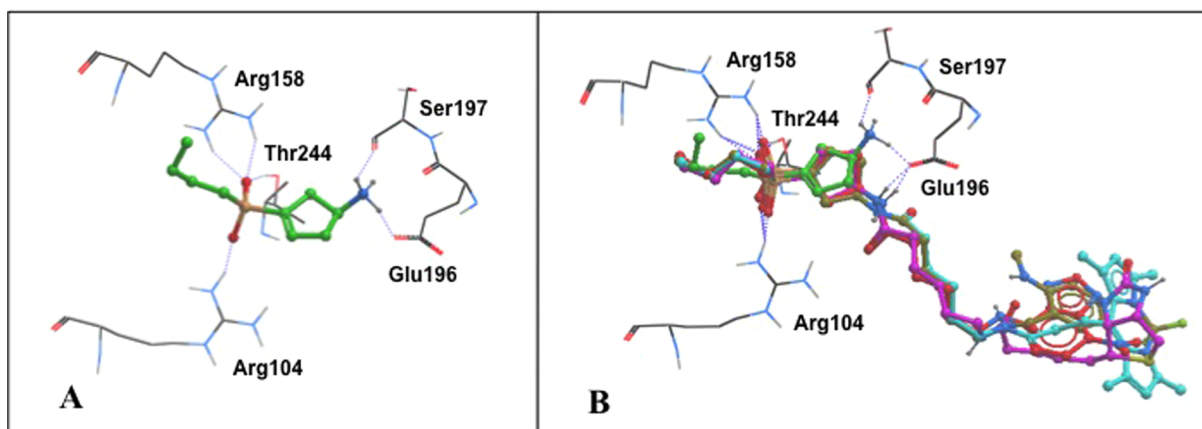


Figure 2. View of the ρ_1 GABA_C ligand-binding site with predicted binding modes. (A) Close-up of (*S*)-4-ACBPBA **6** (green carbons). (B) Close-up of (*S*)-4-ACBPBA-C5-NMA (*S*)-**11d** (brown carbons), (*S*)-4-ACBPBA-C5-NBD **12** (red carbons), (*S*)-4-ACBPBA-C5-BODIPY **13** (aqua blue carbons), and (*S*)-4-ACBPBA-C5-biotin **16** (magenta carbons) that are overlaid on (*S*)-4-ACBPBA **6** in the GABA_C binding site, with the linker and fluorophores external to the orthosteric site.

GABA_C antagonist (*S*)-4-ACBPBA **6** by linking the small fluorophore NMA through linkers of various sizes.

The synthesis of (*S*)-4-ACBPBA-NMA- and (*R*)-4-ACBPBA-C5-NMA-conjugated target compounds (*S*)-**11a–g** and (*R*)-**11d** is depicted in Scheme 1. *N*-Succinimidyl esters **9a–g** were prepared by protection of the amine function of the amino acids with a *tert*-butyloxycarbonyl group (Boc), followed by activation of the acid function of the amino acids with an *N*-hydroxysuccinimide (NHS) (see the Supporting Information for the experimental details). The activated *N*-succinimidyl esters **9a–g** were further reacted with the GABA_C antagonist (*S*)-4-ACBPBA **6** in an aqueous NaHCO₃ solution at room temperature, and subsequent removal of the Boc group using TFA in CH₂Cl₂ to afford compounds **10a–g** in moderate to good yields. Target compounds (*S*)-**11a–g** were obtained by amide coupling of compounds (*S*)-**10a–g** with NHS-activated fluorophore NMA (NMA *N*-succinimidyl ester) in an aqueous NaHCO₃ solution at room temperature. (*R*)-**10d** was obtained from **9d** as described above for the preparation of (*S*)-**10d**. (*R*)-**11d** was synthesized by reaction of (*R*)-4-ACBPBA **7** with the previously prepared ester (*R*)-**10d** under basic conditions at room temperature.

Fluorescent compounds **12** and **13** containing the NBD and BODIPY moieties were prepared as depicted in Scheme 2. Condensation of 7-nitrobenz-2-oxa-1,3-diazol-4-yl chloride (NBD chloride) with (*S*)-**10d** in a K₂CO₃/DMF mixture at room temperature afforded target compound **12** in moderate yield. The synthesis of BODIPY *N*-succinimidyl ester (see the Supporting Information for the experimental details) was achieved in two steps according to a previously described method.²⁸ Compound **13** was obtained in moderate yield by reaction of BODIPY succinimidyl ester and the primary amine of (*S*)-**10d** in a DIPEA/DMF mixture at room temperature.

Biotinylated conjugates **15** and **16** were prepared according to Scheme 3. Formation of an amide bond between biotin *N*-succinimidyl ester **14** and the corresponding primary amine of (*S*)-4-ACBPBA **6** and compound (*S*)-**10d** in a DIPEA/DMF mixture at room temperature afforded biotinylated conjugates **15** and **16**, respectively.

We examined the functional characterization of fluorescent and biotinylated probes on human recombinant GABA receptors expressed in *Xenopus* oocytes using the two-electrode voltage clamp method (Table 1).^{9,10} All 12 probes [(*S*)-**11a–g**,

(*R*)-**11d**, **12**, **13**, **15**, and **16**] were evaluated for activity alone and in the presence of GABA on GABA_A ($\alpha_1\beta_2\gamma_{2L}$), and ρ_1 GABA_C receptors to determine whether they behave as agonists, antagonists, or modulators.

During the initial pharmacological screening, we found that stable and reproducible levels of inhibition of GABA (1 μ M) at ρ_1 GABA_C receptors were reached only after a 5 min preincubation with compounds (*S*)-**11a–g** and (*R*)-**11d** (see the Supporting Information for details). The preincubation needed for some of the compounds for reaching stable and reproducible levels of inhibition of GABA may be due to a molecule's size and lipophilicity needing additional time to diffuse into the binding site. In the (*S*)-4-ACBPBA-NMA series where the NMA group was attached to the primary amine of the ligand through an aminoalkanoyl chain, we observed that compounds with a chain length of fewer than five carbons [compounds (*S*)-**11a–c**] exhibited weak antagonist activity, compared to compound (*S*)-**11d** with a five-carbon chain. The activity for compounds (*S*)-**11e–g** decreased with an increasing chain length. These compounds had a chain length of 6–10 carbons, indicating that a linker of five carbons is optimal. The decrease in activity for compounds (*S*)-**11a** and (*S*)-**11b** (spacer size of fewer than four carbons) could be explained by the relative rigidity caused by the amido-containing chain with the ligand moiety at the binding site. With spacer sizes of more than six carbons, the fluorescent tag may occupy a less favorable area, resulting in a decrease in inhibitory activity.

The optimal chain length of the linker is important for the introduction of molecular flexibility for more convenient positioning of the ligand into the receptor. After optimization of the chain length for fluorescent probes, we linked (*R*)-4-ACBPBA through a 6-aminohexanoyl chain to NMA [(*R*)-**11d**] and evaluated its activity on ρ_1 GABA_C receptors. (*R*)-4-ACBPBA **7** is 12-fold less potent than (*S*)-4-ACBPBA **6** at the ρ_1 GABA_C receptor (Figure 1).⁹ However, (*R*)-4-ACBPBA-C5-NMA (*R*)-**11d** exhibited activity almost equal to that of (*S*)-4-ACBPBA-C5-NMA (*S*)-**11d**, indicating *N*-acylation of the amine removed the differential activity of enantiomers.

We also evaluated NBD and BODIPY (**12** and **13**, respectively) containing probes on both ρ_1 GABA_C and $\alpha_1\beta_2\gamma_{2L}$ GABA_A receptors with and without preincubation (Table 1). (*S*)-4-ACBPBA-C5-NBD **12** exhibited poor potency (IC₅₀ = 281.42 μ M) when tested without preincubation, but its

activity was slightly improved ($IC_{50} = 193.27 \mu M$) with a 5 min preincubation at ρ_1 GABA_C receptors. (S)-4-ACPBPA-C5-BODIPY **13** displayed moderate and improved potency ($IC_{50} = 58.61 \mu M$ with preincubation, and $IC_{50} = 103.14 \mu M$ without preincubation) at ρ_1 GABA_C receptors. The increase in activity for (S)-4-ACPBPA-C5-BODIPY **13** but not for (S)-4-ACPBPA-C5-NMA (S)-**11d** and (S)-4-ACPBPA-C5-NBD **12** indicates that bulkier fluorophores linked by a 6-aminohexanoyl chain are more favorable for activity. These results confirmed the prediction of molecular modeling studies that the large hydrophobic region of the receptor is located adjacent to the binding site of the basic amino group of (S)-4-ACPBPA. As expected, probes (S)-**11d**, (R)-**11d**, **12**, and **13** were found to be inactive (at $300 \mu M$) at $\alpha_1\beta_2\gamma_{2L}$ GABA_A receptors.

We also evaluated biotinylated probes of (S)-4-ACPBPA (compounds **15** and **16**) on the ρ_1 GABA_C receptor. Biotin-coupled probes are suitable for subcellular localization of receptors because biotin forms a tight complex with strept(avidin),²¹ which can be visualized by confocal microscopy using antibodies. Direct attachment of the biotin group to (S)-4-ACPBPA was deleterious for activity, (compound **15** was inactive at $300 \mu M$). However, a linker length of five carbons (compound **16**) is sufficient to restore moderate activity ($IC_{50} = 76.54 \mu M$ with preincubation, and $IC_{50} = 147.92 \mu M$ without preincubation) at ρ_1 GABA_C receptors. These findings represent the first demonstration of electrophysiological activity by *N*-acyl derivatives of selective antagonists (S)-4-ACPBPA **6** and (R)-4-ACPBPA **7**.

To identify structural determinants for the observed activity for the ρ_1 GABA_C receptor, we flexibly docked the structures of (S)-4-ACPBPA **6** and fluorescent and biotinylated probes (S)-4-ACPBPA-C5-NMA (S)-**11d**, (S)-4-ACPBPA-C5-NBD **12**, (S)-4-ACPBPA-C5-BODIPY **13**, and (S)-4-ACPBPA-C5-biotin **16** into the ligand-binding site of a ρ_1 GABA_C homology model.²⁹ Figure 2B shows predicted binding modes for (S)-4-ACPBPA-C5-NMA (S)-**11d**, (S)-4-ACPBPA-C5-NBD **12**, (S)-4-ACPBPA-C5-BODIPY **13**, and (S)-4-ACPBPA-C5-biotin **16**, which show modes of binding for probe-linked (S)-4-ACPBPA similar to that of parent (S)-4-ACPBPA **6** (Figure 2A). The binding affinity is largely ascribed to various electrostatic interactions,^{29–31} including (i) a salt bridge interaction between the phosphinic acid and Arg104 and Arg158,^{29,30} (ii) hydrogen bond contacts of the amine of parent ligand or amide of probes with Glu196,⁹ and (iii) hydrogen bond contacts with Thr244.³¹ The docked model of compounds **6**, (S)-**11d**, **12**, **13**, and **16** (Figure 3) orients the (S)-4-ACPBPA moiety in the active site cavity of the ρ_1 GABA_C receptor (shown with wire mesh) and a flexible spacer between the (S)-4-ACPBPA core and probe where the probe is pointing out of or residing in the noninteractive binding site of the receptors, which cannot be ruled out at this stage. The fluorophores or biotin tags are shown by molecular modeling studies to reside in a pocket away from the orthosteric site or extend into the extracellular space.

In conclusion, we have designed and synthesized for the first time fluorescent and biotinylated probes of selective antagonists for ρ_1 GABA_C receptors. Pharmacological studies show that **12**, **13**, and **16** exhibit activity similar to that of the previously reported muscimol-based probes,^{25,26} with the added advantage that they are selective GABA_C receptors. These results offer new knowledge regarding the binding site and receptor flexibility of GABA_C receptors. The described fluorescent and biotinylated probes will be useful tools for localizing,



Figure 3. View of the (S)-4-ACPBPA moiety (wire mesh) in the active site cavity of the ρ_1 GABA_C receptor model and flexible spacer between the (S)-4-ACPBPA core and probes pointing away from the orthosteric site.

visualizing, and studying the physiopathological processes of GABA_C receptors. Further optimization of probes at amino and phosphinic acid moieties of (S)-4-ACPBPA, in vitro pharmacological evaluation, and fluorescent spectroscopic characterizations of all the probes are underway.

■ ASSOCIATED CONTENT

📄 Supporting Information

Synthetic method, characterization of compounds, pharmacology, and molecular modeling. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Notes

The authors declare no competing financial interest.

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